

## United States Patent and Trademark Office

UNITED STATES DEPARTMENT OF COMMERCE United States Patent and Trademark Office Address: COMMISSIONER FOR PATENTS P.O. Box 1450 Alexandria, Virginia 22313-1450 www.uspto.gov

APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.					
10/668,627	09/23/2003	Kazuko Shinozaki	382.1029DIV2	7894					
23280 7.	590 05/18/2006	EXAM	EXAMINER						
	DAVIDSON & KAPPI AVENUE, 14TH FLOOI	KRUSE, D	KRUSE, DAVID H						
NEW YORK,		i.c	ART UNIT	PAPER NUMBER					
•			1638	- <del>-</del>					
			DATE MAILED: 05/18/2000	5					

Please find below and/or attached an Office communication concerning this application or proceeding.

		Application No.	Applicant(s)						
		10/668,627	SHINOZAKI ET AL.						
	Office Action Summary	Examiner	Art Unit						
•	•	David H. Kruse	1638						
Period fo	The MAILING DATE of this communication app or Reply	ears on the cover sheet with the c	orrespondence address						
WHIC - Exter after - If NO - Failu Any r	ORTENED STATUTORY PERIOD FOR REPLY CHEVER IS LONGER, FROM THE MAILING DATE is not of time may be available under the provisions of 37 CFR 1.13 SIX (6) MONTHS from the mailing date of this communication. Period for reply is specified above, the maximum statutory period were to reply within the set or extended period for reply will, by statute, eply received by the Office later than three months after the mailing and patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tim vill apply and will expire SIX (6) MONTHS from a cause the application to become ABANDONEI	I.  lely filed  the mailing date of this communication.  O (35 U.S.C. § 133).						
Status									
1)□	Responsive to communication(s) filed on								
·	•	action is non-final.							
3)	Since this application is in condition for allowar		secution as to the merits is						
,—	closed in accordance with the practice under E	•							
Dispositi	on of Claims		•						
4)⊠	Claim(s) <u>5-8</u> is/are pending in the application.								
	4a) Of the above claim(s) is/are withdraw	vn from consideration.	·						
_	Claim(s) is/are allowed.								
6)⊠	Claim(s) <u>5-8</u> is/are rejected.								
7)	Claim(s) is/are objected to.								
8)□	Claim(s) are subject to restriction and/o	r election requirement.							
Applicati	on Papers								
9)[	The specification is objected to by the Examine	r.							
10)⊠	The drawing(s) filed on 23 September 2003 is/a	are: a)⊠ accepted or b)⊡ objec	ted to by the Examiner.						
	Applicant may not request that any objection to the	drawing(s) be held in abeyance. See	e 37 CFR 1.85(a).						
	Replacement drawing sheet(s) including the correct	ion is required if the drawing(s) is obj	ected to. See 37 CFR 1.121(d).						
11)	The oath or declaration is objected to by the Ex	aminer. Note the attached Office	Action or form PTO-152.						
Priority ι	ınder 35 U.S.C. § 119								
_	Acknowledgment is made of a claim for foreign ☑ All b)☐ Some * c)☐ None of:	priority under 35 U.S.C. § 119(a)	-(d) or (f).						
	1. Certified copies of the priority documents	s have been received.							
	2. Certified copies of the priority documents	s have been received in Application	on No. <u>09/301,217</u> .						
	$3.\square$ Copies of the certified copies of the prior	ity documents have been receive	ed in this National Stage						
	application from the International Bureau								
* S	see the attached detailed Office action for a list	of the certified copies not receive	d.						
Attachmen	t(s)								
1) Notic	e of References Cited (PTO-892)	4) Interview Summary							
	e of Draftsperson's Patent Drawing Review (PTO-948)	Paper No(s)/Mai	Date						
3) ☑ Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08)  Paper No(s)/Mail Date 9/23/2003.  5) ☑ Notice of Informal Patent Application (PTO-152)  6) ☑ Other: See Continuation Sheet.									

Continuation of Attachment(s) 6). Other: Search Report for SEQ ID NO: 8.

Application/Control Number: 10/668,627 Page 2

Art Unit: 1638

#### **DETAILED ACTION**

#### Claim Rejections - 35 USC § 102

1. The following is a quotation of the appropriate paragraphs of 35 U.S.C. § 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(e) the invention was described in a patent granted on an application for patent by another filed in the United States before the invention thereof by the applicant for patent, or on an international application by another who has fulfilled the requirements of paragraphs (1), (2), and (4) of section 371(c) of this title before the invention thereof by the applicant for patent.

The changes made to 35 U.S.C. § 102(e) by the American Inventors Protection Act of 1999 (AIPA) and the Intellectual Property and High Technology Technical Amendments Act of 2002 do not apply when the reference is a U.S. patent resulting directly or indirectly from an international application filed before November 29, 2000. Therefore, the prior art date of the reference is determined under 35 U.S.C. § 102(e) prior to the amendment by the AIPA (pre-AIPA 35 U.S.C. § 102(e)).

2. Claims 5 and 7 are rejected under 35 U.S.C. § 102(e) as being anticipated by Thomashow *et al* (U.S. Patent 6,417,428 B1, filed 23 November 1998, and claiming priority as a continuation-in-part to U.S. Patent Application 09/017,816, filed 3 February 1998).

Thomashow *et al* disclose an isolated DNA encoding the CBF2 transcription factor of SEQ ID NO: 13 in SEQ ID NO: 12. The amino acid sequence of SEQ ID NO: 13 Thomashow *et al* discloses is identical to SEQ ID NO: 8 disclosed by Applicant. Thomashow *et al* disclose a transgenic plant transformed with an isolated DNA encoding SEQ ID NO: 13 operably linked to a stress responsive promoter in Claim 8. Thomashow *et al* disclose that the promoter would cause higher level of expression to occur at a given environmental condition such as temperature and/or dryness (column 18, lines 27-30). Hence, all of the claim limitations have been previously disclosed by Thomashow *et al*. The disclosure of Thomashow *et al* is supported under 35 USC § 112, first paragraph in the parent application U.S. Patent Application 09/017,816 at

Art Unit: 1638

page 74 and in Figure 14, which is identical to that in the Thomashow *et al* '428 patent.

The Examiner has attached a copy of the Sequence Search for Applicants' SEQ ID NO:

8 to the instant Office action for Applicants' convenience.

### Claim Rejections - 35 USC § 103

- 3. The following is a quotation of 35 U.S.C. § 103(a) which forms the basis for all obviousness rejections set forth in this Office action:
  - (a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.
- 4. Claims 6 and 8 are rejected under 35 U.S.C. § 103(a) as being unpatentable over Thomashow *et al* (U.S. Patent 6,417,428 B1, filed 23 November 1998, and claiming priority as a continuation-in-part to U.S. Patent Application 09/017,816, filed 3 February 1998).

The teachings of Thomashow et al are discussed above.

Thomashow *et al* do not teach the complete DNA sequence of Applicants' SEQ ID NO: 7, only the coding sequence of the encoded protein in SEQ ID NO: 12 of Thomashow *et al*.

The instant claims would have been *prima facie* obvious to one of ordinary skill in the art at the time of Applicant's invention because it is unclear from Applicants' teachings if there is any showing of unexpected results by transforming a plant with a DNA comprising the nucleotide sequence as show in Applicants' SEQ ID NO; 7 as opposed to a DNA comprising the coding sequence, nucleotide sequence as show in Thomashow *et al* in SEQ ID NO: 12.

Application/Control Number: 10/668,627

Art Unit: 1638

5. Claims 5-8 are rejected under 35 U.S.C. § 103(a) as being unpatentable over Liu et al (The Plant Cell August 1998, 10: 1391-1406) in view of Shinwari et al (Biochem. Biophys. Res. Comm. September 1998, 250: 161-170).

This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. § 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR § 1.56 to point out the inventor and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. § 103(c) and potential 35 U.S.C. § 102(e), (f) or (g) prior art under 35 U.S.C. § 103(a).

Applicant cannot rely upon the foreign priority papers to overcome this rejection because a translation of said papers has not been made of record in accordance with 37 CFR § 1.55. See MPEP § 201.15.

Liu *et al* teaches a transgenic plant transformed with a DNA encoding a transcription factor protein consisting of the amino acid sequence as shown in SEQ ID NO: 2 (DREB1A) operably linked to the rd29A promoter, that regulates the transcription of genes located downstream of a stress responsive element, wherein the stress is dehydration stress, low temperature stress or salt stress (see Methods on pages 1402-1405).

Liu et al does not teach a DNA encoding the amino acid sequence of SEQ ID NO: 8 or the DNA sequence of SEQ ID NO: 7.

Art Unit: 1638

Shinwari *et al* teach a DNA comprising the sequence of SEQ ID NO: 7 in Figure 5 on page 164.

It would have been *prima facie* obvious to one of ordinary skill in the art at the time of Applicant's invention to modify the teachings of Liu *et al* to transform a plant with a DNA encoding a DREB transcription factor using the DNA taught by Shinwari *et al*. Shinwari *et al* teach that the DREB1C is more strongly expressed and that transforming a plant would be desirable to make stress tolerant transgenic plants (see pages 168-169). Given the success of Liu *et al*, one of ordinary skill in the art would have had a reasonable expectation of success.

#### **Double Patenting**

6. The nonstatutory double patenting rejection is based on a judicially created doctrine grounded in public policy (a policy reflected in the statute) so as to prevent the unjustified or improper timewise extension of the "right to exclude" granted by a patent and to prevent possible harassment by multiple assignees. A nonstatutory obviousness-type double patenting rejection is appropriate where the conflicting claims are not identical, but at least one examined application claim is not patentably distinct from the reference claim(s) because the examined application claim is either anticipated by, or would have been obvious over, the reference claim(s). See, e.g., *In re Berg*, 140 F.3d 1428, 46 USPQ2d 1226 (Fed. Cir. 1998); *In re Goodman*, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); *In re Longi*, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985); *In re Van Ornum*, 686 F.2d 937, 214 USPQ 761 (CCPA 1982); *In re Vogel*, 422 F.2d 438, 164 USPQ 619 (CCPA 1970); and *In re Thorington*, 418 F.2d 528, 163 USPQ 644 (CCPA 1969).

A timely filed terminal disclaimer in compliance with 37 CFR 1.321(c) or 1.321(d) may be used to overcome an actual or provisional rejection based on a nonstatutory double patenting ground provided the conflicting application or patent either is shown to be commonly owned with this application, or claims an invention made as a result of activities undertaken within the scope of a joint research agreement.

Effective January 1, 1994, a registered attorney or agent of record may sign a terminal disclaimer. A terminal disclaimer signed by the assignee must fully comply with 37 CFR 3.73(b).

Application/Control Number: 10/668,627 Page 6

Art Unit: 1638

7. Claims 5-8 are rejected on the ground of nonstatutory obviousness-type double

patenting as being unpatentable over claims 1-4 of U.S. Patent No. 7,045,355.

Although the conflicting claims are not identical, they are not patentably distinct from

each other because the claimed transformed host cell and vector comprising SEQ ID

NO: 7 renders obvious the transgenic plant of the instant application.

Conclusion

8. No claims are allowed.

9. Any inquiry concerning this communication or earlier communications from the

examiner should be directed to David H. Kruse, Ph.D. whose telephone number is (571)

272-0799. The examiner can normally be reached on Monday to Friday from 8:00 a.m.

to 4:30 p.m.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's

supervisor, Anne Marie Grunberg can be reached at (571) 272-0975. The fax

telephone number for this Group is (571) 273-8300.

Any inquiry of a general nature or relating to the status of this application or

proceeding should be directed to the Group Receptionist whose telephone number is

(571) 272-0547.

DAVID H. KRUSE, PH.D.

PRIMARY EXAMINER

David H. Kruse, Ph.D.

3 May 2006

Application/Control Number: 10/668,627 Page 7

Art Unit: 1638

10. Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to (571) 272-0547.

Patent applicants with problems or questions regarding electronic images that can be viewed in the Patent Application Information Retrieval system (PAIR) can now contact the USPTO's Patent Electronic Business Center (Patent EBC) for assistance. Representatives are available to answer your questions daily from 6 am to midnight (EST). The toll free number is (866) 217-9197. When calling please have your application serial or patent number, the type of document you are having an image problem with, the number of pages and the specific nature of the problem. The Patent Electronic Business Center will notify applicants of the resolution of the problem within 5-7 business days. Applicants can also check PAIR to confirm that the problem has been corrected. The USPTO's Patent Electronic Business Center is a complete service center supporting all patent business on the Internet. The USPTO's PAIR system provides Internet-based access to patent application status and history information. It also enables applicants to view the scanned images of their own application file folder(s) as well as general patent information available to the public.

For all other customer support, please call the USPTO Call Center (UCC) at 800-786-9199.

```
Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OM protein -
         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
         830.5
830.5
830.5
818
818
817
817
795.5
795.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seq length: 0
seq length: 2000000000
                                                                                                                                                                                                                                                                                           982
982
982
982
9837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BLOSUM62
Gapop 10.0 ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-668-627-8
1146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                572060 seqs, 82675679 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      April 26, 2006, 12:23:21 ; Search time 23 Seconds
(without alignments)
776.432 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MNSFSAFSEMFGSDYESPVS......VQWNYNFDVEGDDDVSLWSY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0
100.0
100.0
                                                            885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

885.9

88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBSUED_PATENTS_AA:*

| (cgn2 6/ptcdata/1/iaa/5_COMB.pep:*
| (cgn2 6/ptcdata/1/iaa/f_COMB.pep:*
| (cgn2 6/ptcdata/1/iaa/H_COMB.pep:*
| (cgn2 6/ptcdata/1/iaa/H_COMB.pep:*
| (cgn2 6/ptcdata/1/iaa/RE_COMB.pep:*
| (cgn2 6/ptcdata/1/iaa/RE_COMB.pep:*
| (cgn2 6/ptcdata/1/iaa/RE_COMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
            GenCore version 5.1.7
(c) 1993 - 2006 Biocceleration Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gapext 0.5
   US-09-198-119C-13
US-09-301-66A-8
US-09-301-217-8
US-09-601-802D-13
US-08-949-603-2
US-08-949-580-2
US-08-949-580-2
US-08-950-172A-2
US-08-950-119C-2
US-09-301-66A-6
US-09-301-802D-2
US-09-198-119C-15
US-09-198-119C-9
US-09-198-119C-9
US-09-198-119C-9
US-09-198-119C-9
US-09-601-802D-9
US-09-601-802D-9
US-09-601-802D-93
US-09-601-802D-93
US-09-198-119C-71
US-09-198-119C-71
US-09-198-119C-53
US-09-601-802D-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUMMARIES
Sequence 13, Appl
Sequence 8, Appli
Sequence 8, Appli
Sequence 2, Appli
Sequence 6, Appli
Sequence 6, Appli
Sequence 15, Appli
Sequence 17, Appli
Sequence 17, Appli
Sequence 17, Appli
Sequence 187, Appli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Description
```

45	44	43	42	41	40	υ 9	3 8	37	36	ა 5	34	υ U	32	31	30	29	28
760.5	765.5	765.5	768	768	769.5	769.5	775.5	775.5	775.5	775.5	776.5	776.5	780.5	780.5	785.5	785.5	786.5
		66.8				67.1	67.7	67.7	67.7	67.7	67.8	67.8	68.1	68.1		68.5	
283	280	280	258	258	277	277	277	277	252	252	283	283	255	255	251	251	277
N	N	N	N	N	N	N	N	N	N	N	N	N	ผ	N	N	N	N
US-09-198-119C-63	US-09-601-802D-77	US-09-198-119C-77	US-09-601-802D-85	-1190	US-09-601-802D-57	US-09-198-119C-57	US-09-601-802D-61	-09-198-1190	-09-601-802D-	-09-198-119C	US-09-601-802D-49	US-09-198-119C-49	-09-601-802D	-1190-	-09-601-8020	-09	-802D
Sequence 63,	Sequence //,	1	Sequence 85, Appr	0 00	1.	•	•	Sequence 61,	1,	•		•	α	3 6	3 0	ָ ט פ י	69,

# ALIGNMENTS

```
APPLICANT: Zarka, Daniel

APPLICANT: Jiang, Cai-Zhong

ITILE OF INVENTION: Plant Having Altered Environmental Stress Tole

FILE REFERENCE: 1917.713 Seq List

CURRENT PILING DATE: 1998-11-23

PRIOR APPLICATION NUMBER: US 09/018,233

PRIOR PILING DATE: 1998-02-03

PRIOR APPLICATION NUMBER: US 09/018,233

PRIOR APPLICATION NUMBER: US 09/017,816

PRIOR APPLICATION NUMBER: US 09/017,816

PRIOR APPLICATION NUMBER: US 09/018,235

PRIOR APPLICATION NUMBER: US 09/018,235

PRIOR FILING DATE: 1998-02-03

PRIOR PILING DATE: 1998-02-03

PRIOR PILING DATE: 1998-02-03

PRIOR PILING DATE: 1998-02-03

PRIOR PILING DATE: 1998-02-03

PRIOR APPLICATION NUMBER: US 09/017,575

PRIOR FILING DATE: 1998-02-03

PRIOR APPLICATION NUMBER: US 09/018,227

PRIOR APPLICATION NUMBER: US 09/018,227

PRIOR APPLICATION NUMBER: US 09/018,227

PRIOR PILING DATE: 1998-02-03

PRIOR APPLICATION NUMBER: US 09/018,227

PRIOR APPLICATION NUMBER: US 09/018,227

PRIOR PILING DATE: 1998-02-03

PRIOR APPLICATION NUMBER: US 09/018,234

PRIOR PILING DATE: 1998-02-03

NUMBER OF SEQ ID NOS: 95

SOPTWARE: PARENTA VEL. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1
US-09-198-119C-13
  밁
                                      Ś
                                                                                          밁
                                                                                                                                   S
                                                                                                                                                                                                                                                                              US-09-198-119C-13
                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Pat
SEQ ID NO 13
LENGTH: 216
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1
Patent No.
                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 216; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Stockinger,
APPLICANT: Jaglo-Ottos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                         ORGANISM: Arabidopsis thaliana FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13, Application US/09198119C
5. 6417428
1 MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn Ver. 2.0
                                                                                          MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHFIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stockinger, Eric
Jaglo-Ottosen, Kirsten
Gilmour, Sarah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thomashow, Michael
                                                                                                                                                                                      Conservative
                                                                                                                                                                                    100.0%; Score 1146; DB 2;
100.0%; Pred. No. 2.9e-121;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                    Length 216;
                                                                                                                                                                                           Indels
                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tolerance
                                                                                                                                                                                           Gaps
         120
                                                     120
                                                                                                   60
                                                                                                                                                  60
                                                                                                                                                                                           0
```

```
GENERAL INFORMATION:

APPLICANT: KAZUKO, SHINOZAKI

APPLICANT: Mie, KASUGA

TITLE OF INVENTION: Environmental Stress-tolerant Pla

FILE REFERENCE: 382.1029

CURRENT APPLICATION NUMBER: US/09/301,217

CURRENT FILING DATE: 1999-04-28

PRIOR APPLICATION NUMBER: JP292348/1998

PRIOR APPLICANT

TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
ORGANISM: Arabidopsis thaliana US-09-301-217-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 2
US-09-301-666A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-301-217-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-301-666A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8, Application US/09301666A Patent No. 6495742 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 1146; DB 2; Best Local Similarity 100.0%; Pred. No. 2.9e-121; 0: Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8, Application US/09301217
Patent No. 6670528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 8
LENGTH: 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SHINOZAKI, Kazuko
APPLICANT: KASUGA, Mie
TITLE OF INVENTION: Genes Encoding Plant Transcription factors
FILE REFERENCE: 382.1028
CURRENT APPLICATION NUMBER: US/09/301,666A
CURRENT FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: JP228457/1998
PRIOR APPLICATION STATE: 1998-08-12
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                181 SLIDNMAEGMLLPSPSVQWNYNFDVEGDDDVSLWSY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               181 SLLDNMAEGMLLPSPSVQWNYNFDVEGDDDVSLWSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121 AKEIQKAAAEAALNFQDEMCHMTTDAHGLDMEETLVEAIYTPEQSQDAFYMDEEAMLGMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121 AKEIQKAAAEAALNFQDEMCHMTTDAHGLDMEETLVEAIYTPEQSQDAFYMDEEAMLGMS 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  181 SILDNMAEGMILPSPSVQWNYNPDVEGDDDVSLWSY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           181 SLLDNMAEGMLLPSPSVQWNYNFDVEGDDDVSLWSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    121 AKEIQKAAAEAALNFQDEMCHMTTDAHGLDMEETLVEAIYTPEQSQDAFYMDEEAMLGMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 KWVCELREPNKKTRIWLGTFQTAEMAARAHDVAAIALRGRSACLWFADSAWRLRIPESTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 KWYCELREPNKKTRIWLGTFQTAEWAARAHDVAAIALRGRSACLNFADSAWRLRIPESTC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AKEIQKAAAEAALNFQDEMCHMTTDAHGLDMEETLVEAIYTPEQSQDAFYMDEEAMLGMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 216
                                                                                                                                                                                                                                                                                                                             Plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            180
```

```
APPLICANT: Zarka, Daniel
APPLICANT: Jiang, Cai-Zhong
APPLICANT: Jiang, Cai-Zhong
APPLICANT: Jiang, Cai-Zhong
APPLICANT: Haake, Volker
TITLE OF INVENTION: PLANT HAVING ALTERED ENVIRONMENTAL
TITLE OF INVENTION: STRESS TOLERANCE
FILE REFERENCE: 514442000201/MBI0029
CURRENT APPLICATION NUMBER: US/09/601,802D
CURRENT FILING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 09/018,233
PRIOR APPLICATION NUMBER: 09/018,233
PRIOR FILING DATE: 1998-02-03
PRIOR APPLICATION NUMBER: 09/018,235
PRIOR APPLICATION NUMBER: 09/018,235
PRIOR APPLICATION NUMBER: 09/017,575
PRIOR APPLICATION NUMBER: 09/017,575
PRIOR APPLICATION NUMBER: 09/017,575
PRIOR APPLICATION NUMBER: 09/018,237
PRIOR APPLICATION NUMBER: 09/018,237
PRIOR FILING DATE: 1998-02-03
PRIOR APPLICATION NUMBER: 09/018,237
PRIOR FILING DATE: 1998-02-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                                       Ś
                                                                                              밁
                                                                                                                                        á
                                                                                                                                                                                                                                                                                          ; LENGTH: 216
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-09-601-802D-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 216
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1998-11-23
NUMBER OF SEQ ID NOS: 259
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13, Application US/09601802D Patent No. 6706866
                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1998-02-03
PRIOR APPLICATION NUMBER: 09/198,119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     181 SILDNMAEGMILPSPSVQWNYNFDVEGDDDVSIWSY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            121 AKEIQKAAAEAALNFQDEMCHMTTDAHGLDMEETLVEAIYTPEQSQDAFYMDEEAMLGMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 AKEIQKAAAEAALNFQDEMCHMTTDAHGLDMEETLVEAIYTPEQSQDAFYMDEEAMLGMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           181 SLLDNMAEGMLLPSPSVQWNYNFDVEGDDDVSLWSY 216
61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 KWVCELREPNKKTRIWLGTFQTAEMAARAHDVAAIALRGRSACLNFADSAWRLRIPESTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 KWYCELREPNKKTRIWLGTFQTAEMAARAHDVAAIALRGRSACLNFADSAWRLRIPESTC
                                               61 KWVCELREPNKKTRIWLGTFQTAEMAARAHDVAAIALRGRSACLNFADSAWRLRIPESTC 120
                                                                                                                                             1 MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                MNSFSAFSEMFGSDYESPVSSGGDYSPKLATSCPKKPAGRKKFRETRHPIYRGVRQRNSG
KWVCELREPNKKTRIWLGTFQTAEMAARAHDVAAIALRGRSACLNFADSAWRLRIPESTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gilmour, Sarah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stockinger, Eric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jaglo-Ottosen,
                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                1998-11-23
OS: 259
                                                                                                                                                                                         100.0%; Score 1146; DB 2; Length 100.0%; Pred. No. 2.9e-121; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 1146; DB 2;
100.0%; Pred. No. 2.9e-121;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kirsten
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                   216;
                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 120
  120
                                                                                                   60
                                                                                                                                                  60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                 0
```